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(54) Title: IM PEPTIDES

#### (57) Abstract

The invention is directed to the diagnosis and treatment of herpes virus related diseases. By noting within the Epstein-Barr virus (EBV) open reading frames (ORFs) transcribed late in the viral cycle for which a translation product may or may not have been established, and synthesizing one or more polypeptides each of which includes at least one segment, each segment comprising at least part of the amino acid sequence identified in that ORF, a specific and reliable diagnostic test for, and treatment of, infectious mononucleosis and related diseases is possible. The preferred amino acid sequences for these segments include NSPKNG, KNGSNQ, SNQLVI, AHARDK, RDKAGA, VMAMIL, SEPRPR and PSRTPS. These sequences can be further combined to provide polypeptides with at least one segment comprising sequences selected from AHARDKAGAVMAMIL, ASLNSPKNGSNQLVI, ELESEPRPRPSRTPS, QAMKKIEDKVRKSVD, SRSRGREAKKVQISD, LIKASLRKDRKLYAE, VSFSKTRRAIRESRA, CNYSAGEEDDQYHAN, RPHRRPVSKRPTHKP, EITQEENRGEQRLGH, GALRARLDRPRPTAQ, NSPKNGSNQAHARDKSEPRPR, NSPKNGSNQRDKAGASEPRPR, NSPKNGSNQSEPRPRKNGSNQ, NSPKNGSNQLVISEPRPRPSRTPS, NSPKNGSNQLVIPSRTPS and NSPKNGSNQAHARDKAGASEPRPR.

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#### TITLE: IM PEPTIDES

### TECHNICAL FIELD

THIS INVENTION is directed to the diagnosis and treatment of herpes virus related diseases. In particular, it relates to the use of specific open reading frames (ORFs) within the Epstein-Barr virus (EBV) which encode antigens recognized by EBV-specific antibodies raised during infectious mononucleosis (IM) and related diseases and the use of synthetic peptides based on the amino acid sequences encoded by these ORFs in a specific and reliable diagnostic test for, and treatment of, IM and related diseases.

#### BACKGROUND ART

- Epstein-Barr virus (EBV) is a member of the herpes virus family and is present in all human populations. Primary infection usually occurs in early childhood and remains silent throughout a person's life. However, when uninfected adolescents and young adults are exposed to EBV, about 60% manifest infectious mononucleosis (IM).
- The predominant laboratory test used to establish the diagnosis of IM has been the demonstration of heterophil antibodies. The rapid slide tests have become the most widely used method to detect these heterophil antibodies. In contrast, quantitative agglutination tests, such as the Paul-Bunnell-Davidsohn method, are more accurate but are also more tedious and time consuming.

The use of tests to measure heterophil antibodies have limitations, namely, only between 80-95% of IM patients produce these antibodies, these antibodies are absent in

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a large percentage of young children and the antibodies are produced in a variety of other diseases such as lymphoma, hepatitis and leukemia. The measurement of heterophil antibodies also does not give any indication of the severity of the disease and cannot be used to monitor the course of IM.

Immunofluoresence tests that measure antibodies to EBV can also be used in the diagnosis of heterophilnegative cases of IM, or patients with atypical manifestations.

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These tests, however, are time consuming and require the use of trained personnel and specialized equipment which does not make them amenable to the routine analysis of large numbers of samples.

- The diagnosis of an acute primary EBV infection can also be determined by an IgM response to EBV-viral capsid antigen (VCA). The VCA is composed of a large number of different antigens. Components of VCA are defined by the fact that they are expressed late in the replicative cycle of the virus. Many VCA components have been mapped to specific open reading frames (ORF's) within the EBV genome though there are many ORF's, known to be expressed late in replication, to which specific VCA antigens have not yet been identified.
- synthetic polypeptide 25 Genetic engineering and technologies now enable the manufacture of quantities of protein and polypeptide antigens. However, these techniques are only effective if the amino acid residue sequence of the native protein is 30 known.

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The amino acid residue sequence of a natural protein can be determined by sequencing of the protein itself.

Alternatively, the DNA sequence that codes for the protein may also reveal the protein's amino acid residue sequence.

Antibodies can be used to determine whether an ORF present in a DNA sequence codes for a protein. involves manufacturing an array of protein fragments or whose amino acid polypeptides residue 10 sequences correspond to the hypothetical sequences obtained from the ORFs. The protein fragments or polypeptides to which naturally occurring antibodies immunoreact thereby identify the ORF as encoding a The complete amino acid naturally occurring protein. 15 sequence of this protein could then be deduced from the DNA sequence of the ORF.

#### DISCLOSURE OF THE INVENTION

It is a general object of the present invention to overcome, or at least ameliorate, one or more of the 20 above disadvantages, and to provide a specific and reliable test for the diagnosis for, and treatment of, IM and related diseases.

As the complete DNA sequence for EBV has been identified, including the start and stop codons which, 25 prima facie, define potential ORFs for the transcription of the genetic code, the present inventors have synthesized peptides based on the predicted amino acid sequences encoded by these ORFs, even though the proteins to which these ORFs may relate were not first 30 established as being produced by the virus, and have

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demonstrated that EBV-specific antibodies raised during IM react with the synthesized peptides.

Thus, according to a first aspect of the present invention, there is provided a peptide comprising a sequence which includes at least one segment which codes for an antigen recognized by EBV-specific antibodies raised during IM or a related disease.

As a second aspect, the present invention also includes within its scope a method of identifying a polypeptide 10 suitable for use in the diagnosis of IM and related diseases, said method comprising:

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- (1) noting within EBV open reading frames transcribed late in the viral cycle for which a translation product may or may not have been established;
- (2) synthesising one or more polypeptides each of which includes at least one segment wherein each segment comprises at least part of the amino acid sequence identified in that open reading frame; and
  - (3) determining whether said polypeptide is effective in the diagnosis of IM and related diseases.
- 25 When sera from patients who exhibited clinical symptoms of IM and related diseases were assayed using the peptides of the invention, a positive reaction was noted with a high correlation between this assay and known assays for identifying IM and related diseases.

5

Therefore, according to a third aspect of the present invention, there is provided a method of diagnosis of infectious mononucleosis or a related disease, said method comprising assaying serum from a patient suspected of having infectious mononucleosis or a related disease with at least one peptide as hereinbefore defined.

The present invention also provides, as a fourth aspect, a kit for use in the diagnosis of IM or a 10 related disease, said kit comprising:

- (a) at least one peptide as hereinbefore defined; and
- (b) a means for indicating the presence of a reaction, particularly an immunoreaction, between said at least one peptide and another molecule(s), especially anti-EBV antibodies.

Since a patient with IM or a related disease contains antibodies to the peptides of the invention, it is likely that the peptides of the invention, when administered to the patient, would elicit anti-EBV antibodies.

Therefore, as a fifth aspect of the present invention, there is provided a vaccine that, when administered, is capable of inducing antibodies effective against EBV, said vaccine comprising:

- (a) at least one peptide as hereinbefore defined; and
- (b) a carrier and/or diluent and/or adjuvant.

6

As used throughout the specification, the term "carrier or diluent" denotes an organic or inorganic, natural or synthetic material with which the active ingredient is combined in order to facilitate the administration of 5 the vaccine of the invention. This carrier or diluent therefore, generally inert and it must Similarly, pharmaceutically acceptable. the "adjuvant" has the usual meaning in the art to describe a material which aids the operation of the active 10 ingredient.

According to a sixth aspect of the present invention, there are also provided antibodies and substantially whole antibodies raised to - or induced by - the peptides of the invention as hereinbefore defined.

15 These molecules are collectively referred to as receptors and can be raised in animal hosts using the vaccine as hereinbefore defined.

Preferably, the peptide of the invention comprises at least one segment selected from the following 20 sequences:

AHARDKAGAVMAMIL
ASLNSPKNGSNQLVI
ELESEPRPRPSRTPS

25 QAMKKIEDKVRKSVD
SRSRGREAKKVQISD
LIKASLRKDRKLYAE
VSFSKTRRAIRESRA
CNYSAGEEDDQYHAN

30 RPHRRPVSKRPTHKP
EITQEENRGEQRLGH
GALRARLDRPRPTAQ

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More preferably, the peptide of the invention comprises at least one segment selected from the following sequences:

5 NSPKNG

KNGSNQ

SNQLVI

AHARDK

RDKAGA

10 VMAMIL

SEPRPR

**PSRTPS** 

Most preferably, the peptide of the invention comprises at least one segment selected from the following 15 sequences:

NSPKNGSNQAHARDKSEPRPR NSPKNGSNQRDKAGASEPRPR NSPKNGSNQSEPRPRKNGSNQ NSPKNGSNQLVISEPRPRPSRTPS

20 NSPKNGSNQLVIPSRTPS

NSPKNGSNQAHARDKAGASEPRPR

A particularly preferred peptide of the invention comprises at least one segment containing the sequence:

#### NSPKNGSNQLVIPSRTPS

25 All amino acid residues identified throughout the specification are in the natural or L-configuration. In

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keeping with standard polypeptide nomenclature, abbreviations for amino acid residues are as follows:

	SYMBOL	AMINO ACID
	Y	L-tyrosine
5	G	glycine
	F	L-phenylalanine
	M	L-methionine
	A	L-alanine
	s	L-serine
10 .	I	L-isoleucine
	L	L-leucine
	T	L-threonine
	v	L-valine
	P	L-proline
15	К	L-lysine
	н	L-histidine
	Q	L-glutamine
	E	L-glutamic acid
	W	L-tryptophan
20	R	L-arginine
	D	L-aspartic acid
	N	L-asparagine
	c .	L-cysteine

#### DETAILED DESCRIPTION OF EMBODIMENTS

#### 25 SUBJECTS, MATERIALS AND METHODS

### Preparation And Use Of Synthetic Peptides

Peptides (15 aa each) were synthesized by the Multiple Simultaneous Peptide technique (MSPS) of Houghton, R.A. (1985, Proc. Natl. Acad: Sci. USA 82, 5131-5135). The synthetic peptides were linked to bovine serum albumin (BSA) with glutaraldehyde as described by Bulinski et

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al. (1983, Proc. Natl. Acad. Sci. USA 80, 1506-1510).
Essentially, 5mg of dry peptide were added to 0.5ml
(4mg/ml) BSA in 100mM phosphate buffer, pH 7.3. To this
were added 0.25ml of 0.25% (v/v) glutaraldehyde for each
5 mg of dry peptide. The solution was left in the dark
over night at 21°C to conjugate, after which the
solution was extensively dialyzed against PBS containing
50mM glycine, pH 7.3. The conjugated peptides were then
stored at -20°C until needed. Peptide sequences,
10 deduced from 13 different ORFs transcribed late in
viral replication, were synthesized. These peptides are
identified in Table 1.

10

# TABLE 1 VCA PEPTIDES

	ORF	PEPTIDES	NUMBER
	BCRF1	AENQDPEAKDHVNSL	1
5	BCRF1	FFQTKDEVDNLLLKE	7
	BDLF2	AKAEERTAEMDDTMA	3
	BDLF2	GGMKRKQCRVDRLTD	8
	BDLF3	AHARDKAGAVMAMIL	2
	BDLF3	PTVPDERQPSLSYGL	12
10	BKRF2	ASLNSPKNGSNQLVI	4
	BILF2	CNYSAGEEDDQYHAN	20
	BILF2	RPHRRPVSKRPTHKP	23 -
	BSRF1	EEPETFECPDRWRAE	. 5
	BGLF1	EITQEENRGEQRLGH	21
15	BGLF1	VSFSKTRRAIRESRA	18
	BALF1	LIKASLRKDRKLYAE	9
	BALF1	YAVFTRDEKDLPLPA	19
	BBRF3	ELESEPRPRPSRTPS	6
	BBRF3	RSSTSSSSSRSTRRQ	15
20	BXRF1	GALRARLDRPRPTAQ	22
	BXRF1	PRSARAGRAGGRKGQ	11
	BORF1	MKVQGSVDRRRLQRR	10
	BORF1	RGSEFTRDVRGLVEE	14
	BLRF2	QAMKKIEDKVRKSVD	13
25	BLRF2	SRSRGREAKKVQISD	16
	BMRF2	TSGLERRRSIFCARG	17

#### ELISA assay

The stock peptide-BSA conjugate was diluted 1/100 with DD H<sub>2</sub>O and 50 µl of the peptide-BSA conjugate were added each well ïn 96 well microtiter plates (Flow 5 Laboratories) and the plates were left overnight at 37°C To block the wells 200µl of a solution containing 5% BSA, 0.5M carbonate buffer pH 9.0 were added and left for 30 minutes at ambient temperature. The wells were then washed four times with 200  $\mu$ l of 0.1% BSA-PBS/1%Tween. Human sera (diluted 1 in 100 in 5%BSA/2xPBS/1%Tween) were added and incubated at 21°C The plates were then washed six times with for 1h. 0.1%BSA/2xPBS/0.1%Tween and 100 ul of peroxidaselabelled anti-human IgM (Tago, µ fraction) (diluted 1/5000 in 5%BSA/2xPBS/0.1%Tween) were added and then the plates incubated at 37°C for 30 minutes. The plates were again washed six times as above. The plates were given a rinse with distilled water (neutral pH), and substrate 1 m M ABTS (2,2-azino-bis(3-(100  $\mu l$ οf ethlybenzthiazoline-6-sulphonic acid) diammonium salt 20 St. Louis, USA) in 100mM phosphate-citrate (Sigma, (pH 4.3) containing 0.004% (v/v) hydrogen buffer peroxide) were added and the plates incubated at 37°C for 30 minutes. Finally the plates were read at 410nm.

#### 25 Subjects and sera

30

Samples of sera (26) were obtained from patients with IM. These patients were diagnosed as having clinical symptoms of IM and were confirmed by immunofluoresence assays for IgM and IgG antibodies to VCA and by immunoblotting for antibodies to early antigens (EA) and absence of antibodies to EBV nuclear antigen 1 (EBNA1). Sera (14 were EBV seropositive and 8 EBV seronegative) were also obtained from healthy controls and their EBV

status was determined by immunofluoresence assays to the EBV antigens. The normal controls were all negative for IgM antibodies to EBV. For the clinical trials, hundreds of samples of serum, from patients displaying IM-like symptoms, were collected.

#### Immunofluoresence assay for KBV antigens.

Anti-VCA titres were measured according to the method of Henle and Henle. (J Bacteriol. 91, 1248-1256).

#### Heterophil antibody assay

10 The Paul Bunnell test was used to measure heterophil antibody titres in serum.

#### RESULTS AND DISCUSSION

#### Screening of EBV peptides

The 23 synthetic peptides were initially screened against 26 samples of sera from confirmed IM patients, 8 EBV seronegative controls and 14 EBV seropositive controls. The results, presented in Fig. 1, illustrate that a number of peptides were reactive with IgM antibodies from IM patients while showing little reaction with the sera from either EBV seropositive or seronegative controls. Peptides # 2, 4, 6, 13, 16 and 18 were selected for further studies.

## Reaction of IM and normal sera with selected KBV peptides.

25 Comparison of total Ig and IgM reactions with the peptides indicated that normal seropositive individuals lacked both IgM and IgG antibodies to these peptides

However, it appeared that IM patients (Fig. 2). contained only IgM antibodies to these peptides. Measurement of IgG antibodies to the peptides, in both IM sera and sera from normal controls, confirmed the 5 absence of these antibodies (results not shown).

These results indicated that measurement of either IgM or total Ig could be useful in identifying serum samples from IM patients.

Since only IqM antibodies to the peptides appear to be 10 present in IM patients, there should not be a problem with rheumatoid factor or interference from antibodies, both of which are usually a problem with indirect ELISA assays.

#### Clinical study

- In order to ascertain the viability of using the 15 peptides to identify cases of IM, sera were obtained from patients who showed clinical symptoms of These sera were assayed by ELISA, using the 6 selected synthetic peptides, by immunofluoresence presence of IgM antibodies to VCA and for heterophil 20 antibodies. Patients were considered to have IM if they were both heterophil positive (titres of 1/16 or higher) IgM positive (titres of 1/40 or higher) immunofluorescence or heterophil negative but with positive cells and IgM 25 atypical monnuclear immunofluoresence. The ELISA based IgM assay using the synthetic peptides was considered to be positive when a serum reacted with four or more of the peptides (A410 of 0.40 or above). The results of the assays, obtained
- with all of the patients, are shown in Table 2. 30

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TABLE 2

		TABLE	5 2		
	Patient	Heterophil	ATLs (%)	ELISA (IgM)	IFA (IgM)
	1	<10	0	-	<10
5	2	64	13	+	160
	3	<10	0	-	<10
	4	<10	0	-	<10
	5	<10	0	-	<10
	<u>6</u>	<10 .	0	-	<10
10	1 2 3 4 5 6 7	<10	0	-	<10
		<10	0	<del>-</del>	<10
	9	<10	0	+	160
	10	<10	0	-	<10
1 =	11	<10	0	-	<10
15	12 · 13	<10 <10	0 0	<b>-</b>	<10
	14	<10	0	_	<10 <10
	15	<10	0	_	<10
	16	<10	0	_	<10
20	17	<10	ŏ	_	<10
20	18	<10	ŏ		<10
	19	<10	Ö	_	<10 .
	20	<10	0	_	<10
	21	<10	0	_	<10
25	22	<10	15	+	160
	23	<10	0	_	<10
	24	256	21	+	160
	25	<10	0		<10
	26	<10	0	_	<10
30	27	<10	0	-	<10
	28	<10	0		<10
	29	<10	14	-	<10
	30	<10 <10	3 0	_	<10
35	31 32	<10	0	. <u>-</u>	<10 <10
35	33 .	10	0	_	160
	34	<10	Ŏ	_	<10
	35	<10	ŏ	_	<10
	36	<10	ŏ	_	<10
40	37	<10	Ö	_	<10
••	38	<10	0 0 0	_	<10
	39	<10	Ö	_	<10
	40	<10	24	-	<10
	41	<10		_	<10
45	42	<10	0	-	<10
	43	<10	0 .	-	<10
	44	<10	16	-	<10
	45	<10	0	-	<10
_	46	64 .	24	+	360
50	47	<10	0 0 2 0	-	<10
	48	16	0	-	<10
	49	<10	2	-	<10
	50	<10	U	_	<10

15

TABLE 2 (cont)

	Patient	Heterophil	ATLs (%)	ELISA (IgM)	IFA (IgM)
5	51 52 53	<10 <10 <10	0 0 3 0	- -	<10 <10 10
10	54 55 56 57	<10 <10 64 <10	65 0	- - -	<10 <10 320 <10
	58 59 60	<10 <10 <10	0 0 0	- - -	<10 <10 <10
15	61 62 63 64	<10 <10 16 256	0 1 0 39	- - +	<10 <10 <10 160
20	65 66 67	<10 <10 <10	9 0 0	- -	<10 <10 <10
	68 69 70 71	<10 <10 <10 <10	0 0 0	- - - +	<10 <10 <10 <10
25	72 73 74	256 <10 <10	23 0 0 9	+	320 <10 <10
30	75 76 77	<10 <10 <10 <10	9 0 0 0	+ -	40 <10 <10 <10
	78 79 80 81	<10 <10 <10 <10	0	- - -	<10 <10 <10 <10
35	82 83 84	<10 <10 <10	0 0 0	- -	<10 <10 <10
40	85 86 87 88	<10 64 <10 64	0 19 . 0 4	- + - +	<10 160 <10 160
	89 90 91	<10 <10 64	4 0 0 45	- - +	<10 <10 160
45	92 93 94	<10 <10 <10 256	0 0 0 32	- - - +	<10 <10 <10 160
50	95 96 97 98 99	<10 <10 <10 <10	0 0 0 0	- - -	<10 <10 <10 <10

30

16

TABLE 2 (cont)

	Patient	Heterophil	ATLs (%)	ELISA (IgM)	IFA (IgM)
	100	<10	3	_	<10
5	101	<10	0	_	<10
	102	<10	0	_	<10
	103	<10	0	-	<10
	104	16	30	+	160

results of the statistical analysis of the 10 correlation between the three assays systems are presented in Table 3.

TABLE 3 CORRELATIONS AND POLYCHORIC TEST STATISTICS

•		ELISA	IF(IgM)	HETEROPHIL
15	ELISA IF(IgM) HETEROPHIL	1.000	0.988 1.000	0.901 0.964 1.000

#### ASYMPTOTIC VARIANCES OF ESTIMATED CORRELATIONS

	ELISA-IF(IgM)	=	0.00020
20	ELISA-HETEROPHIL	=	0.00407
	IF(IgM)-HETEROPHIL	=	0.00094

The results obtained from the statistical analysis demonstrate a 96% correlation between the IF-IgM assay and the heterophil antibody assay which better the 25 reported data. However, the correlation between the ELISA assay and IF-IgM was 99% indicating that this assay was more accurate at predicting patients with IM. There was one serum which was ELISA positive but IF-IgM negative and it is possible that this patient may have not as yet developed a high enough titre of IgM antibodies to be detected by the IF-IgM test. Two sera samples were positive by IF-IgM and negative by the ELISA assay.

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Peptides # 2,4 and 6 were selected for further clinical trials, whereby 510 samples of serum, from patients displaying IM-like symptoms, were assayed for heterophil antibody using the Paul Bunnell test and for the presence of IgM antibodies to EBV by immunofluoresence as described above. These results were then correlated with the results obtained using the ELISA assay with the three synthetic EBV peptides and with the Monolert assay as marketed by Johnson and Johnson. The statistical analysis of these results are shown below.

#### CORRELATIONS AND TEST STATISTICS (PC=POLYCHORIC)

			•	TEST OF	MODEL	TEST OF	ZERO CORR.
	Corre	elat.	Chi-	squ.DF	P-VALUE	CHI-SQU.	P-VALUE
	IFA VS. PB	.868	(PC)	.000 0	1.000	889.303	.000
15	ELISA VS. PB	.882	(PC)	.000 0	1.000	969.017	.000
	ELISA VS. IFA	.897	(PC)	.000 0	1.000	1073.819	.000
	MONO VS. PB	.718	(PC)	.000 0	1.000	413.573	.000
	MONO VS. IFA	.595	(PC)	.000 0	1.000	238.506	.000
	MONO VS. ELISA	.614	(PC)	.000 0	1.000	259.295	.000

20 ESTIMATED CORRELATION MATRIX

		PB	IFA	ELISA	MONO
			<del></del>		
	PB	1.000			
	IFA	0.868	1.000		
25	ELISA	0.882	0.897	1.000	
	MONO	0.718	0.595	0.614	1.000

PB - Paul Bunnell

IFA - Immunofluoresence assay for EBV-specific IgM

ELISA - ELISA assay using the synthetic EBV peptides of the invention

30 MONO - Monolert assay as marketed by Johnson and Johnson

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These results show a much higher correlation between the synthetic peptides of the invention and the IFA assay than did the PB test or the Monolert assay.

Rather than use three independent peptides in the assay system, longer peptides containing combinations of B-cell epitopes (ie the epitopes to which the antibodies bind) from each of the 15 amino acid peptides were constructed. To define the B-cell epitopes in each of the three peptides overlapping 6 amino acid synthetic peptides were prepared and assayed by ELISA using a pooled IM positive sera and a pooled normal sera.

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5

#### ORIGINAL 15-mers PEPTIDES

#4 ASLNSPKNGSNQLVI

**#2 AHARDKAGAVMAMIL** 

#6 ELESEPRPRPSRTPS

#### SYNTHETIC OVERLAPPING PEPTIDES

		PEPTIDE	POSITIVE SERA	NEGATIVE SERA
10		ASLNSPKNGSNQL	VI	
		ASLNSP	0.287	0.110
	**	NSPKNG	0.555	0.143
	**	KNGSNQ	0.537	0.141
	**	SNQLVI	0.437	0.133
15		AHARDKAGAVMAM	T.	
	**	AHARDK	0.639	0.215
	**	RDKAGA	0.496	0.122
		AGAVMA	0.324	0.099
	**	VMAMIL	0.505	0.170
20		ELESEPRPRPSRTI	PS .	
		ELESEP	0.150	. 0.063
	**	SEPRPR	0.623	0.228
		RPRPSR	0.407	0.131
	**	PSRTPS	0.546	0.185
				_

<sup>25 \*\* -</sup> INDICATES THE PEPTIDES CONTAINING B-CELL EPITOPES.

20

Having defined the B-cell epitopes contained within the original 15 aa peptides, different combinations of these epitopes were used in the synthesis of larger synthetic peptides. Examples of some of these combinations are shown below:

#### EPITOPE COMBINATIONS: .

	PEPTIDE	A.	NSPKNGSNQAHARDKSEPRPR
	PEPTIDE :	в.	NSPKNGSNQRDKAGASEPRPR
-	PEPTIDE	c.	NSPKNGSNQSEPRPRKNGSNQ
10	PEPTIDE :	D.	NSPKNGSNQLVISEPRPRPSRTPS
	PEPTIDE :	E.	NSPKNGSNQLVIPSRTPS
	PEPTIDE :	F.	NSPKNGSNQAHARDKAGASEPRPR

These 6 peptides were then tested for their reaction with sera from known IM patients and with sera from 5 normal controls. The sera were all well characterized for the presence/absence of IgM antibodies to EBV by immunofluoresence. The results of that testing are presented in Table 4.

TABLE 4

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_													
Peptide	A *	B **	IM1	IMZ	IM3	IM4	IM5	NOR1	NOR9	NOB3	NOR1 NOR2 NOR3 NOR4 ACM	MODE	200M
									71011	MOIN	NON	NORS	NOKe
А	1.58	0.25	1.34	1.34	1.34 1.34 1.52 1.53 0.63	1.52	1.53	0.63	0.16	0.16 0.31 0.13	0 12	26 0	0.47
										70.0	. 1	0.40	0.40
В	1.64	0.19	1.20	1.35	1.35   1.24   1.63   1.15   0.48	1.63	1.15		0,19	0.19 0.95 0.11		06 0	96.0
											٦		0.50
೮	1.04	0.12	0.81	1.11	1.11   0.93   1.47   0.85   0.21	1.47	0.85		ر بر	0 15 0 39 0 19		1,4	70
									7:0	70.0		0.14 0.24	0.24
Д	1.31	0.17	0.97	1.14	1.14   1.05   1.49   0.87   0.36	1.49	0.87		1.0	76	10		
								1	01.0	0.64	17.0 0.14 0.12 0.14	0.14	17.0
田	1.20	0.12	0.64	1.07	0.88	1.46	0.67	1.07 0.88 1.46 0.67 0.95	r r	0 15 0 17	-		
								9.50	01.0	0.I (		0.13 0.21	0.21
ᄄ	1.55	0.03	1.15	1.22	1.19	1,60	1 19	0 27	. 91	06.0	1.22 1.19 1.60 1.19 0.37 0.91 0.90 0.15		
									77.0	60.0	CT:0	27.0	

\* Pooled control IM serum

\*\* Pooled control normal serum
IM Infectious mononucleosis
NOR Normal

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These results demonstrate that any of these epitope combinations could be used to identify serum from IM patients. However some of the peptides gave reasonably strong reactions with sera from normal individuals (peptides A,B and F in particular). Of those peptides which had low reactions with the normal sera peptide E gave the lowest values and so was chosen for further studies.

Peptide E was assayed with a larger contingent of sera from patients with IM and normal controls. Control wells, containing only BSA, were also included in the assay to check for sera which may react with the carrier protein. The results of that assay are presented in Table 5.

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TABLE 5
ELISA ASSAY OF PEPTIDE E.

		ELISA ASSAY OF		
	SERA			WELL (BSA)
	A	Positive control	0.993	0.066
5	В	Negative control	0.116	0.068
	1	IM	1.050	0.077
	2	IM	1.133	0.072
	3	IM	0.689	0.056
	4	IM	0.780	0.064
10	5	IM	0.852	0.085
	6	IM	0.598	0.051
	7	IM	0.813	0.053
	8	IM	0.953	0.062
	9	IM	1.093	0.051
15	10	IM	0.739	0.054
	11	IM	1.128	0.043
	12	IM	0.933	0.056
	13	MI	0.795	0.058
	14	IM	0.556	0.051
20	15	IM	0.717	0.047
	16	IM	1.454	0.065
	17	Normal	0.101	0.049
	18	Normal	0.187	0.055
	19	Normal	0.098	0.051
25	20	Normal	0.072	0.052
	21	Normal	0.149	0.045
	22	Normal	0.061	0.050
	23	Normal	0.073	0.044
	24	Normal	0.080	0.045
30	25	Normal	0.120	0.047
	26	Normal	0.159	0.052
	27	Normal	0.142	0.057
	28	Normal	0.054	0.046
	29	Normal	0.270	0.076
35	30	Normal	0.336	0.045
	31	Normal	0.280	0.051
	32	Normal	0.337	0.041
	33	Normal	0.136	0.042

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These results demonstrate that peptide E could be used to reliably detect cases of IM while showing little reaction with sera from normal individuals.

- The clinical and other data obtained indicates that the ELISA assay using the peptides of the present invention is a specific and reliable test for the diagnosis of infectious mononucleosis and related diseases. The present invention should also find use in the treatment of such diseases.
- 10 Those skilled in the art will appreciate that the above embodiments are given by way of exemplification of the invention only, and that changes may be made to the details set out therein without departing from the scope of the invention as defined in the following claims.

#### **CLAIMS**

- A peptide comprising a sequence which includes at least one segment which codes for an antigen recognized by EBV-specific antibodies raised during IM or a related disease.
  - A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence AHARDKAGAVMAMIL.
- 3. A peptide as defined in Claim 1, wherein said at 10 least one segment comprises the sequence ASLNSPKNGSNQLVI.
  - 4. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence ELESEPRPRPSRTPS.
- 15 5. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence QAMKKIEDKVRKSVD.
- 6. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence LIKASLRKDRKLYAE.
  - 7. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence VSFSKTRRAIRESRA.
- 8. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence CNYSAGEEDDQYHAN.
  - 9. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence RPHRRPVSKRPTHKP.

- 10. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence EITQEENRGEQRLGH.
- 11. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence GALRARLDRPRPTAQ.
  - 12. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence SRSRGREAKKVQISD.
- 10 13. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNG.
  - 14. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence KNGSNQ.
- 15. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence SNQLVI.
  - 16. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence AHARDK.
  - 17. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence RDKAGA.
- 20 18. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence VMAMIL.
  - 19. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence SEPRPR.
- 20. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence PSRTPS.

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- 21. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNGSNOAHARDKSEPRPR.
- 22. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNGSNQRDKAGASEPRPR.
  - 23. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNGSNOSEPRPRKNGSNO.
- 10 24. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNGSNQLVISEPRPRPSRTPS.
  - 25. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNGSNOLVIPSRTPS.
    - 26. A peptide as defined in Claim 1, wherein said at least one segment comprises the sequence NSPKNGSNOAHARDKAGASEPRPR.
- 27. A peptide as defined in any one of Claims 1 to 26 20 in association with any suitable carrier and/or diluent and/or adjuvant.
  - 28. A method of identifying a polypeptide suitable for use in the diagnosis of IM and related diseases, said method comprising:
- 25 (1) noting within EBV open reading frames transcribed late in the viral cycle for which a translation product may or may not have been established;

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- (2) synthesising one or more polypeptides each of which includes at least one segment wherein each segment comprises at least part of the amino acid sequence identified in that open reading frame; and
  - (3) determining whether said polypeptide is effective in the diagnosis of IM and related diseases.
- 10 29. A method of diagnosis of infectious mononucleosis or a related disease, said method comprising assaying serum from a patient suspected of having infectious mononucleosis or a related disease with at least one peptide as defined in any one of Claims 1 to 27.
  - 30. A kit for use in the diagnosis of IM or a related disease, said kit comprising:
    - (a) at least one peptide as defined in any one of Claims 1 to 27; and
- 20 (b) a means for indicating the presence of an immunoreaction between said peptide and anti-EBV antibodies.
  - 31. A kit for use in the diagnosis of IM or a related disease, said kit comprising:
- 25 (a) at least one peptide as defined in any one of Claims 1 to 27; and
  - (b) a means for indicating the presence of a reaction between said peptide and another molecule.

- 32. A vaccine that, when administered, is capable of inducing antibodies effective against infection by EBV, said vaccine comprising:
- (a) at least one peptide as defined in any one of Claims 1 to 26; and
  - (b) a carrier and/or diluent and/or adjuvant.
  - 33. Antibodies and substantially whole antibodies raised to or induced by at least one peptide as defined in any one of Claims 1 to 27.
- 10 34. A peptide as defined in Claim 1 substantially as described with reference to the Examples.
  - 35. A method as defined in Claim 28 substantially as described with reference to the Examples.

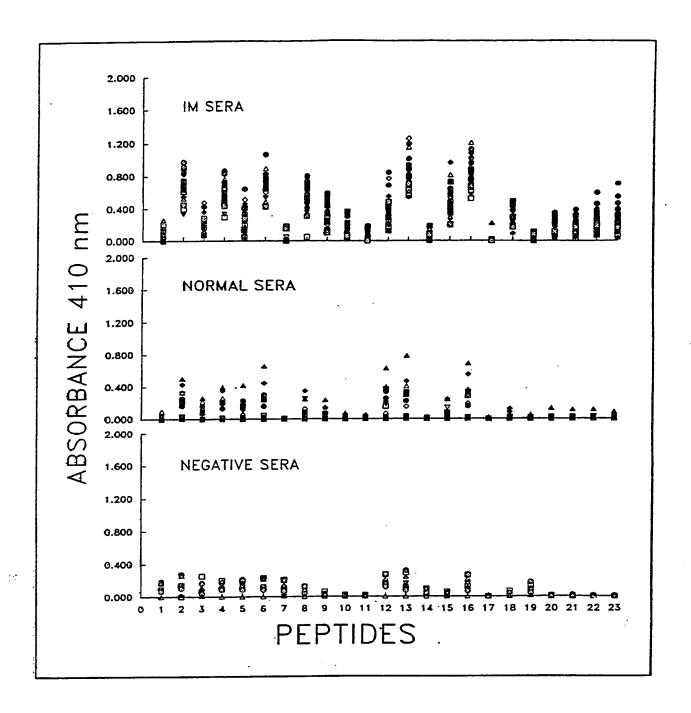


FIGURE 1

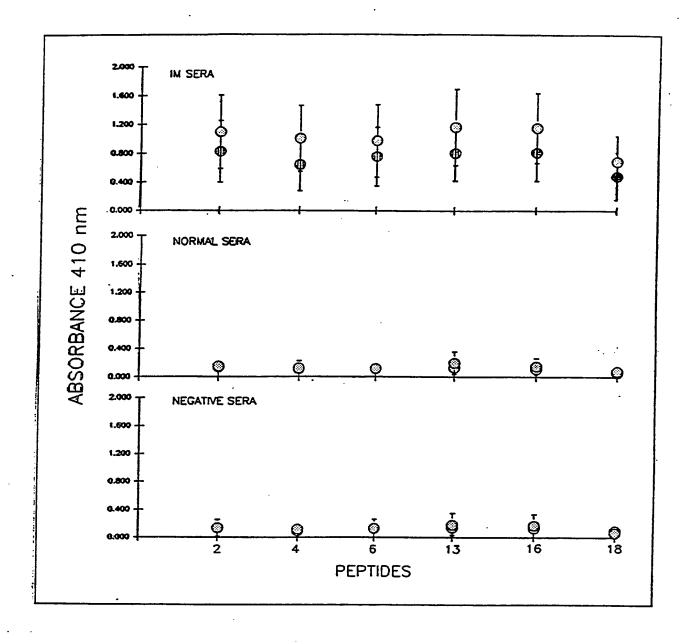


FIGURE 2

## INTERNATIONAL SEARCH REPORT

International Application No. PCT/AU 90/00564

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		F SUBJECT MATTER (if several c				
According to International Patent Classification (IPC) or to both National Classification and IPC						
Int. Cl.	<sup>5</sup> COTK 7/06,	7/08, 7/10, 15/12, C12N 7/00,	C12Q 1/70, A61K 39/245, G01	N 33/569		
II. FIE	LDS SEARCHED					
		Hini	mum Documentation Searched 7			
Classific	ation System	Classific	ation Symbols			
AU: IPC	5	C07K 7/06, 7/08, 7/10, C0	7C 103/52, GO1N 33/569			
<u> </u>		cumentation Searched other tha ent that such Documents are In		d 8		
Chem. A STN and	bs. KEYWORDS PIR databases	Epstein () Barr () Virus OR s for Int. searching of seqs.	EBV and IM OR Infectious Monor of claims 13-20.	onucleosis.		
III. DOC	UMENTS CONSIDE	CRED TO BE RELEVANT 9				
Category*	Citation	of Document, With indicati of the relevant passag		Relevant to   Claim No 13		
X	AU,A, 82073,	/87 (SCRIPPS CLINIC AND RESEAR see pages 5,6 and claims	CH FOUNDATION) 9 June 1988	1 only		
X,P		/89 (SCRIPPS CLINIC AND RESEAR .90) see page 13 and claims	CH FOUNDATION) 22 February	1 only		
   X 	US 4707358 ( and claims	(KIEFF et al) 17 November 1987	(17.11.87) see example 4	1 only		
x   x 	Nature, Volu   "DNA sequence   see whole do	ume 310, issued 1984 July, (Lo ce and expression of the B95-8 ocument	ndon) R. Baer et al, Epstein-Barr virus genome"	1 only		
	1    -		(continued)	†   		
* Spec	 cial categorie	s of cited documents: 10 *T	<ul> <li>later document published international filing dat</li> </ul>			
art   part   "E" earl	which is not ticular releva lier document		and not in conflict with cited to understand the underlying the invention X document of particular re claimed invention cannot	the application but principle or theory elevance; the		
"L" docu cla publ othe	ument which ma im(s) or which lication date er special rea	y throw doubts on priority is cited to establish the of another citation or " son (as specified)	or cannot be considered inventive step  Y" document of particular reclaimed invention cannot	elevance; the be considered to		
"O" document referring to an oral disclosure, involve an inventive step when the documer use, exhibition or other means is combined with one or more other such document published prior to the documents, such combination being obvious						
	ernational fil priority date	ing date but later than eclaimed	a person skilled in the a document member of the sa			
IV. CER	TIFICATION					
		letion of the	Date of Mailing of th	is International		
	onal Search		Search Report	1		
i	ry 1991 (27.02		1 March 1991			
]	onal Searching		Signature by Jutiforty	ed Officer  CEDRIC SCHAFFER		
Anctrolis	n Patent Offic	92	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	CELECUS SAIRETER		

X ·	Journal of Virological Methods, Volume 21, issued 1988 (The Netherlands) J. Middeldorp and P. Herbrink "Epstein-Barr virus specific marker molecules for early diagnosis of infectious mononucleosis", see whole document	1 cally
x	Nucleic Acids Research, Volume 16 no.7, issued 1988 (IRL Press Limited, Oxford, England) D. Walls et al "The analysis of EBV proteins which are antigenic in vivo", see whole document	1 only
A	EP,A, 0316170 (G GENE GALWAY LIMITED) 17 May 1989 (17.05.89)	

V. [] ODSERVALIDAD WHERE CERTAIN CLAIMS WERE FOOD CHARACTERISTS.

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

- 1.[] Claim numbers ..., because they relate to subject matter not required to be searched by this Authority, namely:
- 2.[] Claim numbers , because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
- 3.[] Claim numbers ..., because they are dependent claims and are not drafted in accordance with the second and third sentences of PCT Rule 6.4 (a):

#### VI. [ ] OBSERVATIONS WHERE UNITY OF INVENTION IS LACKING 2

This International Searching Authority found multiple inventions in this international application as follows:

- 1.[] As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims of the international application.
- 1 2.[ ] As only some of the required additional search fees were timely paid by the applicant, this
  international search report covers only those claims of the international application for
  which fees were paid, specifically claims:
  - 3.[] No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claim numbers:
- 4. [] As all searchable claims could be searched without effort justifying an additional fee, the International Searching Authority did not invite payment of any additional fee.

! Remark on Protest

- [ ] The additional search fees were accompanied by applicant's protest.
- [ ] No protest accompanied the payment of additional search fees.

## ANNEX TO THE INTERNATIONAL SEARCH REPORT ON INTERNATIONAL APPLICATION NO. PCT/AU 90/00564

This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

	ent Document ed in Search Report			Paten	t Family Mem	bers	·
US	4707358	EP	151079	JР	60232094	-	
EP	316170	JP	2002399				
UA	44000/89	WO	9001495				
AU	82073/87	DK JP ZA	6388/87 63225398 8709025	EP NZ	280813 222794	IL US	84690 487921 <b>3</b>

END OF ANNEX